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February 16, 2003, 16:42:40; Search time 2267.72 Seconds (without alignments) 15566.316 Million cell updates/sec
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1: CgnZ_6/ptodata/I/pna/US06_COMB.seq:*

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10: /cgn2_6/ptodata/I/pna/US09_BCO
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ptodata/1/pna/US100A_COMB.seq:*
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries

    nucleic search, using sw model

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45: /cgn2_6/ptodata/1/pna/US6001_COMB.seq:*
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84: /cgn2_6/ptodata/1/
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Description	Section 2			3.	œ	44	74	7.5	, ,			10	14	, 6, 6	, ,		2 0		, c	2 0	Sequence 86, Appl	
ID	US-09-497-967-5	US-09-497-967-102	US-09-497-967-53	US-09-497-967-3	US-09-498-612-8	US-09-497-967-44	US-09-497-967-74	US-09-497-967-75	US-09-497-967-70	US-09-497-967-71	US-09-497-967-72	US-09-497-967-79	US-09-497-967-76	US-09-497-967-73	US-09-497-967-77	US-09-497-967-82	US-09-497-967-78	US-09-497-967-84	US-09-497-967-83	US-09-497-967-85	US-09-497-967-86	
DB	18	18	18	18	18	18	18	18	18	18	18	18	18	18	1.8	18	18	18	18	18	18	
Query Match Length DB	1404	1410	1404	1404	1404	1410	138	123	117	104	100	100	66	95	95	95	94	94	92	92	92	
ouery Match	100.0	100.0	8.66	55.7	55.7	55.7	9.8	8.8	7.5	7.4	7.1	7.1	7.1	6.8	6.8	6.8	6.7	6.7	9.9	9.9	9.9	
Score	1404	1404	1400.8	782.6	782.6	782.6	138	123	105	104	100	100	66	95	95	95	94	94	92	92	92	
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13409, A
11300, A
859, App
                         Sequence 1, Appli
Sequence 7, Appli
Sequence 2, Appli
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GRENKAL INFORMATION:
APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Lin, Tian-Long
ITILE OF INVENTION: DICHEVAPOPHTHINUS
TITLE OF INVENTION: ICHTHYOPHTHINUS
TITLE OF INVENTION: ICHTHYOPHTHINUS
TITLE OF INVENTION: ICHTHYOPHTHINUS
TITLE REPRENCE: 235.00170101
CURRENT APPLICATION NUMBER: 60/131,121
PRIOR APPLICATION NUMBER: 60/131,121
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-02-04
PRIOR FILING DATE: 1999-03-03-03
PRIOR PROLICATION NUMBER: 60/122,372
PRIOR PELLING DATE: 1999-03-03-03
PRIOR PELLING DATE: 1999-03-03-03
PRIOR PELLING DATE: 1999-03-17
PRIOR PELLING DATE: 1999-03-17
SPRIOR PELLING DATE: 1999-03-17
SPRIOR PELLING DATE: 1999-03-17
SOFFWARE: PALENTIN Ver. 2.1
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18 US-09-497-967-81

18 US-09-497-967-80

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18 US-09-498-612-7

18 US-09-498-612-7

19 US-09-498-612-4

10 US-09-498-612-4

10 US-09-498-612-4

10 US-09-498-612-4

10 US-09-196-161D-9

10 US-07-763-352A-14

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10 US-09-196-161-1

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10 US-09-115-5

10 US-09-115-5

10 US-09-115-5

10 US-09-116-1425

10 US-09-186-1425

10 US-09-186-1425

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10 US-10 US-11300

US-10 US-113100

US-10 US-138-13100

US-60-173-469-859
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100.0%; Pred. No. 0;
Live 0; Mismatches
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     Best Local Similarity 100.
Matches 1404; Conservative
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APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
TITLE OF INVENTION: ICHTHYOPHTHIRUS
TITLE OF INVENTION: ICHTHYOPHTHIRUS
FILE REPERENCE: 235.0170101
CURRENT APPLICATION NUMBER: 60/9497,967
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-02-04
PRIOR PAPLICATION NUMBER: 60/118,634
PRIOR RELING DATE: 1999-03-04
PRIOR PAPLICATION NUMBER: 60/122,372
PRIOR FILING DATE: 1999-03-07
PRIOR PELING DATE: 1999-03-07
PRIOR FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 102
SOFTWARE: Patentin Ver: 2.1
                                                        CCTGAGTCTGCTAAGAAGAACATCCAGTGTGACTTCGCTAACTTCCTGTCTATCTCTG 1380
                                                                  GCTAACIGICCIGIGGAACCGAGACCAACACGCTGGACAGGIGGACGACCTGGGAACC 120
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100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                       CTGCTGATCTCTTACTACCTGCTG 1404
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; GENERAL INFORMATION:
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Matches 1404; Conservative
                                                                                                                                          RESULT 2
US-09-497-967-102
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US-09-497-967-102
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                         APPLICANT: Clark, Theodore G.
APPLICANT: Clark, Theodore G.
APPLICANT: Clark, Theodore G.
APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
TITLE OF INVENTION: LOTHWOPHTHINUS
TITLE OF INVENTION: OLOTOOLO
TURRENT FILING DATE: 2000-02-04
CURRENT FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-02-04
PRIOR FILING DATE: 1999-03-02
PRIOR FILING DATE: 1999-03-02
PRIOR FILING DATE: 1999-03-07
SPRIOR FILING DATE: 1999-03-07
SPRIOR FILING DATE: 1999-03-17
SOFTWARE FILING DATE: 1999-03-17
SOFTWARE PAPLICATION NUMBER: 60/124,905
PRIOR FILING DATE: 1999-03-17
SOFTWARE PATENTION NUMBER: 60/124,905
SPRIOR FILING DATE: 1999-03-17
SOFTWARE PATENTION NUMBER: 60/124,905
SOFTWARE PATENTIANT NUMBER: 60/124,905
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Pred. No. 0;
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         Sequence 53, Application US/09497967 GENERAL INFORMATION:
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Matches 1402; Conserv
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OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                  Q ID NO 53
LENGTH: 1404
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Sequence Application US/09497967
Sequence Application US/09497967
Sequence Application:
APPLICANT: Clark, Theodore G.
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
TITLE OF INVENTION: CUTHYOPHTHIRUS
FILE REFERENCE: 235.00170101
CURRENT APPLICATION NUMBER: US/09/497,967
CURRENT FILING DATE: 2000-02-04
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                                      AACACCCCTTTCAACCCTGGAAAGTCTCAGTGTACCCCTTGTCCTGCTATCAAGCCTGCT
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PRIOR APPLICATION NUMBER: 60/131,121
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US-09-497-967-3
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Sequence 8, Application US/09498612
Sequence 8, Application US/09498612
Sequence 8, Application US/09498612
Sequence 8, Application:
APPLICANT: GAERFIG, Jacek
APPLICANT: GLARK, Theodore G.
APPLICANT: CLARK, Theodore G.
APPLICANT: CLARK, Theodore G.
APPLICANT: TITLE OF INVENTION: RECOMBINAT EXPRESSION OF HETEROLOGOUS NUCLEIC ACIDS IN TITLE OF INVENTION: PROTOZOA
FILE REFERENCE: 235.00100101
CURRENT APPLICATION NUMBER: 05/09/498,612
CURRENT APPLICATION NUMBER: 60/122,372
PRIOR FILING DATE: 1999-03-07
PRIOR FILING DATE: 1999-03-07
PRIOR FILING DATE: 1999-03-17
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1090-02-04
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                                                                                               GCTGCTACCCTGGCTAAGCAGTGTAACATCGCTTGTCCTGACGGAACCGCTATCGTTCT
                                                                                                               ; TYPE: DNA
; ORGANISM: Ichthyophthirius multifillis
US-09-498-612-8
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1381 TTATTGATTTCTTATTA 1397
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Pred. No. 9.6e-210;
0; Mismatches 384; Indels
PRIOR FILING DATE: 1999-04-27
PRIOR PLING DATE: 1999-02-04
PRIOR FILING DATE: 1999-02-04
PRIOR FILING DATE: 1999-02-04
PRIOR PRICATION NUMBER: 60/122,372
PRIOR PRILING DATE: 1999-03-02-02
PRIOR PRILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 102
SOFTWARE: PACENTIN VET: 2.1
SEQ ID NO 3
                                                                                                                                                                ; ORGANISM: Ichthyophthirius multifiliis US-09-497-967-3
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Best Local Similarity 72.5%;
Matches 1013; Conservative
                                                                                                                                                       TYPE: DNA
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Length 1404;
               Indels
 DB 18;
Score 782.6; DB 18;
Pred. No. 9.6e-210;
0; Mismatches 384;
  55.7%;
72.5%;
                 Conservative
           Similarity
                 Matches 1013;
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Pred. No. 9.6e-210;
0; Mismatches 384;
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Matches 1013; Conservative
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LENGTH: 1410
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US-09-497-967-44
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US-09-497-967-44
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APPLICANT: Clark, Theodore G.
APPLICANT: Clark, Theodore G.
APPLICANT: Clark, Theodore G.
APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
TITLE OF INVENTION: ICHTHYOPHTHIRIUS
FILE REFERENCE: 235.001701.
CURRENT APPLICATION NUMBER: U5/09/497,967
PRIOR APPLICATION NUMBER: G0/131,121
PRIOR APPLICATION NUMBER: 60/131,121
PRIOR PLING DATE: 1999-04-27
PRIOR PLING DATE: 1999-04-27
PRIOR PLING DATE: 1999-03-04
PRIOR PLILNG DATE: 1999-03-03-04
PRIOR PLILNG DATE: 1999-03-03-07
PRIOR PLILNG DATE: 1999-03-03-17
NUMBER OF SEQ ID NOS: 102
SOFTWARE: Patentin Ver. 2.1
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1321 CCTGAGTCTGCTAAGAAGAACATCCAGTGTGACTTCGCTAACTTCCTGTCTATCTCTCTG 1380
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Sequence 75 Application US/09497967

Sequence 75 Application US/09497967

GENERAL INFORMATION:

APPLICANT: Clark, Theodore G.

APPLICANT: Lin, Than Long

TITLE OF INVENTION: ICHTHYOPHTHIRIUS

TITLE OF INVENTION: ICHTHYOPHTHIRIUS

FILE REFERENCE: 235.0017010

CURRENT APPLICANTION NUMBER: US/09/497,967

PRIOR APPLICATION NUMBER: 60/131,121
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OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: oligonucleotide primers
US-09-497-967-7-0
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100.0%; Pred. No. 5.3e-28;
ative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                    ; Sequence 74, Application US/09497967
; GENERAL INFORMATION:
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                                                                                                                                                     1381 CTGCTGATCTCTTACTA 1397
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Matches 138; Conservative
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RESULT 10
US-09-497-967-71/C
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APPLICANT: Clark, Theodore G.
APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
TITLE OF INVENTION: ICHTHYOPHTHINGS
TITLE OF INVENTION: ICHTHYOPHTHINGS
FILE REPERENCE: 235.00170101
CURRENT APPLICATION NUMBER: US/09/497,967
CURRENT FILING DATE: 1000-02-04
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-03-02-04
PRIOR FILING DATE: 1999-03-02
PRIOR FILING DATE: 1999-03-02
PRIOR FILING DATE: 1999-03-02
PRIOR FILING DATE: 1999-03-02
PRIOR FILING DATE: 1999-03-07
SEQ ID NOS: 102
SCFWARE: Datentin Ver. 2.1
SEQ ID NO 70
LENGTH: 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   490 GCTIGICCIACCGGAACCGCICIGGACGACGGAGIGACCACCGACIACGIGCGCICITIC 549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 ATGAAGAACAACATCCTGGTGATCCTGATCTCTCTTCTTCATCAACCAGATCAAGTCT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              430 CGCGTGGGAGGAGCTCTGACCGCTGGAAACGCTGCTACCATCGTGGCTCAGTGTAACGTG 489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13 ATGAAGAACAACATCCTGGTGATCCTGATCTCTCTGTTCATCAACCAGATCAAGTCT 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     63 GCTTGTCCTACCGGAACCGCTCTGGACGACGAGTGACCACCGACTACGTGCGCTCTTTC 4
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                                                                                                                                                                                                                                                                                                                                                                                         8.8%; Score 123; DB 18; Length 123; 100.0%; Pred. No. 8.7e-24; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 7.5%; Score 105; DB 18; Length 1 Best Local Similarity 100.0%; Pred. No. 1e-18; Matches 105; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: oligonucleotide primers US-09-497-967-70
                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: oligonucleotide primers US-09-497-967-75
PRIOR FILING DATE: 1999-04-27
PRIOR APPLICATION NUMBER: 60/118,634
PRIOR FILING DATE: 1999-02-04
PRIOR APPLICATION NUMBER: 60/122,372
PRIOR FILING DATE: 1999-03-02
PRIOR FILING DATE: 1999-03-07
PRIOR FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 70, Application US/09497967; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 123; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 550 ACC 552
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                                                                                                                                                                                                         SEQ ID NO 75
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GENERAL INCORATION:
GENERAL INCORATION:
APPLICANT: Clark Theodore G.
APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: DIGMOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
TITLE OF INVENTION: ICHTHYOPHTHINUS
FILE REFERENCE: 235.00170101
CURRENT FILING DATE: 236.00170101
CURRENT FILING DATE: 1999-04-27
PRIOR APPLICATION NUMBER: 60/131,121
PRIOR APPLICATION NUMBER: 60/131,121
PRIOR APPLICATION NUMBER: 60/122,372
PRIOR PRILING DATE: 1999-04-27
PRIOR PELLING DATE: 1999-03-04
PRIOR FILING DATE: 1999-03-02
PRIOR FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 102
SOFTWARE: PALENTIN VET: 2.1
                                                                                                                                                                                                                                                                                               APPLICANT: Clark, Theodore G.
APPLICANT: Clark, Theodore G.
APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
TITLE OF INVENTION: ICHHHYOPHTHINUS
FILE REPERENCE: 235.001701J
CURRENT APPLICATION NUMBER: US/09/497,967
CURRENT APPLICATION NUMBER: 06/131,121
PRIOR APPLICATION NUMBER: 60/131,121
PRIOR APPLICATION NUMBER: 60/131,237
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-03-04
PRIOR FILING DATE: 1999-03-04
PRIOR FILING DATE: 1999-03-05
PRIOR FILING DATE: 1999-03-05
PRIOR FILING DATE: 1999-03-17
SPRIOR FILING DATE: 1999-03-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84 GACCAACACCGCTGGACAGGTGGACGACCTGGGAACCCCTGCTAACTGTGTGAACTGTCA 143
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                               144 GAAGAACTICTACTACAACAACGCTGCTGCTTTCGTGCCTGGAG 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence: ; OTHER INFORMATION: oligonucleotide primers US-09-497-967-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 GCTAACTGTCCTGTGGGAACCGAGACCAACACGCTGGACAGGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
7.4%: Score 104; DB 18;
Best Local Similarity 100.0%; Pred. No. 1.9e-18;
Matches 104; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                    ; Sequence 71, Application US/09497967; GENERAL INFORMATION:
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242 CTCCTGCTACCGCTAACCTGGTGACCCAGTGTAACGTGAAGTGTCCTGCTGGAACGCTA 301
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US-09-497-967-73/c
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APPLICANT: Clark, Theodore G.
APPLICANT: Dickerson, Jr., Harry W.
APPLICANT: Lin, Than-Long
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
TITLE OF INVENTION: ICHTHYOPHTHIRIUS
FILE REFERENCE: 235.0017011
CURRENT APPLICATION NUMBER: 05/131,121
PRIOR APPLICATION NUMBER: 06/131,121
PRIOR APPLICATION NUMBER: 60/131,121
PRIOR APPLICATION NUMBER: 60/131,372
PRIOR PELING DATE: 1999-04-27
PRIOR APPLICATION NUMBER: 60/122,372
PRIOR PELING DATE: 1999-03-04
PRIOR FILING DATE: 1999-03-03-02
PRIOR FILING DATE: 1999-03-03-07
PRIOR FILING DATE: 1999-03-03-17
NUMBER OF SEQ ID NOS: 102
SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                    165 CGCTGCTGCTTTCGTGCCTGGAGCTTCTACCTGTACCCCTTGTCCTCAGAAGAAGAAGG
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GRNERAL INFORMATION:
GRNERAL INFORMATION:
GRNERAL INFORMATION:
GRNERAL INFORMATION:
APPLICANT: Clark Theodore G.
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                              753 GAACAACTGGGTGGCTCAGAACACCGAGTGTACCAACTGTGCTCCTAACTTCTACAACAA 812
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                                                                                                                                                                     Length 100;
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                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                           OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: oligonucleotide primers US-09-497-967-72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Oligonucleotide primers US-09-497-967-79
                                                                                                                                                                                                                                                                                                                7.1%; Score 100; DB 18;
100.0%; Pred. No. 2.5e-17;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      813 CAACGCTCCTAACTTCAACCCTGGAAACTCTACCTGTCTG 852
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US-09-497-967-79/c
; Sequence 79, Application US/09497967
; GENERAL INFORMATION:
                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 79
LENGTH: 100
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                       Query Match
Best Local Similarity 100.0
Matches 100; Conservative
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Matches 100; Conservative
LENGTH: 100
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICAMT: OLARK, Theodoce G.
APPLICAMT: Dickerson, Jr., Harry W.
APPLICAMT: Lin, Tian-Long
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES OF
TITLE OF INVENTION: DIAGNOSTIC AND PROTECTIVE ANTIGEN GENE SEQUENCES
TITLE OF INVENTION: Lin, Tian-Long
TITLE OF INVENTION: LINGUIDER: 0.50/0.02-04
FILE REFERENCE: 235.001710101
CURRENT APPLICATION NUMBER: 60/131,121
PRIOR APPLICATION NUMBER: 60/131,121
PRIOR APPLICATION NUMBER: 60/132,372
PRIOR APPLICATION NUMBER: 60/122,372
PRIOR PELING DATE: 1999-02-04
PRIOR PELING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 102
SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 7.1%; Score 99; DB 18; Length 99; Best Local Similarity 100.0%; Pred. No. 4.9e-17; Matches 99; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 95;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: oligonucleotide primers
US-09-497-967-76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence:
CTHER INFORMATION: Oligonucleotide primers
US-09-497-967-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                592 AACAACGGAAACACCCCTTTCAACCCTGGAAAGTCTCAG 630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 6.8%; Score 95; DB 18; L
Best Local Similarity 100.0%; Pred. No. 6.5e-16;
Matches 95; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: ICHTHYOPHTHIRIUS
CURRENT APPLICATION NUMBER: US/09/497,967
CURRENT FILING DATE: 2000-02-04
PRIOR PELICATION NUMBER: 60/131,121
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-02-04
PRIOR FILING DATE: 1999-02-04
PRIOR APPLICATION NUMBER: 60/118,634
PRIOR FILING DATE: 1999-03-02
PRIOR APPLICATION NUMBER: 60/122,372
PRIOR PILING DATE: 1999-03-02
PRIOR APPLICATION NUMBER: 60/124,905
PRIOR APPLICATION NUMBER: 60/124,905
PRIOR PRILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 102
SOFTWARE: PALENT VET: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
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RESULT 15
US-09-497-967-77/C
Sequence 77, Application US/09497967
Sequence 77, Application US/09497967
SEQUENCE INFORMATION:
APPLICANT: Clark, Theodore G.
APPLICANT: Lin, Tian-Long
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: LIHATY W.
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: LIHATY W.
APPLICANT: Lin, Tian-Long
TITLE OF INVENTION: LIHATY WHYOURTHIRIUS
TITLE OF INVENTION: LIHATY W.
CURRENT APPLICATION NUMBER: US/09/497,967
CURRENT FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-04-27
PRIOR FILING DATE: 1999-02-04
PRIOR FILING DATE: 1999-03-07
PRIOR FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 102
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 77
LEBORTE. THE SECOND NOT THE SEQ ID NOS: 102
LEBORTE. PATENTER P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: oligonucleotide primers US-09-497-967-77
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